<u>**In the Description**</u>

Under 37 C.F.R. § 1.121(b), please amend the specification as follows:

On page 1, after the title of the invention, please add the following section headings and accompanying paragraph:

CROSS-REFERENCE TO RELATED APPLICATIONS

This application is a U.S. national application under 37 C.F.R. § 371(b) of International Application Serial No. PCT/GB2005/001030 filed March 17, 2005, which claims the priority benefit of United Kingdom application serial no. 0406054.7 filed March 17, 2004.

BACKGROUND AND SUMMARY OF THE INVENTION

On page 3, line 7, please amend as follows:

(i)   ILLWQPIPV (PAP.135) ~~SEQ. ID.~~ SEQ ID NO: 1,

On page 3, line 17, please amend as follows:

(i)   CPRFQELESETLKSE (PAP.161) ~~SEQ. ID.~~ SEQ ID NO: 2,

On page 12, line 23, please insert the following heading:

BRIEF DESCRIPTION OF THE DRAWINGS

On page 14, line 17, please insert the following heading:

DETAILED DESCRIPTION

On page 14, please amend the partial paragraph beginning on line 20 as follows:
Candidate peptides with either HLA-A2*0201 or HLA-DRB1*0401/HLA-DRB1*0101 binding motifs were identified using the SYFPEITHI on-line epitope prediction algorithm , which analyses peptides for the presence of certain amino acid residues which favour MHC binding . The peptide corresponding to positions 58-66 (GILGFVFT - SEQ ID NO: 13) of the influenza virus M1 protein has been previously identified as a potent HLA-A2*0201 CTL epitope and was employed as a positive control in CTL generation assays. For class-II proliferation assays, the influenza peptide corresponding to positions 307-319 of the influenza virus (PKYVKQNTLKLAT - ~~SEQ. ID.~~ SEQ ID NO: 3) was used. The peptide corresponding to positions 128-140 (TPPAYRPPNAPIL - ~~SEQ. ID.~~ SEQ ID NO: 4) of the hepatitis-B pre-core protein (AAK57285) is a known mouse MHC class-II

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On page 21, line 10, please delete the Table and replace it with the following:

Seq. ID. No.	Amino-acid sequence	Name	Description
SEQ ID NO: 5	ALDVYNGLL	PAP.299	
SEQ ID NO: 6	VLAKELKFV	PAP.30	
SEQ ID NO: 7	IMYSAHDTTV	PAP.284	novel (predicted) epitope
SEQ ID NO: 8	ILLWQPIPV	PAP.135	novel (predicted) epitope
SEQ ID NO: 9	ALASCFCFFC	PAP.15	novel (predicted) epitope
SEQ ID NO: 10	PQGFGQLTQLGMEQH	PAP.64	novel (predicted) epitope
SEQ ID NO: 11	CPRFQELESETLKSE	PAP.161	novel (predicted) epitope
SEQ ID NO: 12	SKVYDPLYSESVHNF	PAP.207	novel (predicted) epitope

On page 21, line 24, please amend as follows:

peptide corresponding to positions 58-66 (GILGFVFT - ~~SEQ. ID.~~ SEQ ID NO: 13) of the influenza

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